Amendments to the Claims:

The following listing of claims will replace all prior versions and listings of claims in the application.

Listing of Claims:

- 1. (Original) A method of producing a DNA array, characterized by comprising the steps of:
- (1) preparing a mixture of DNA fragments in which a modified base or a base is exposed,
- (2) bringing the mixture of DNA fragments obtained in step (1) into contact with an antibody specific to the modified base or the base, and separating the mixture into a group consisting of DNA fragments which form an immunocomplex with the antibody and another group consisting of DNA fragments which do not react with the antibody, or a group consisting of DNA fragments showing a high affinity for the antibody and another group consisting of DNA fragments showing a low affinity for the antibody,
- (3) identifying all or part of DNA fragments contained in each of the DNA fragment groups, and
- (4) arranging one or more nucleic acids capable of hybridizing with any one of the identified DNA fragments on a substrate.
- 2. (Original) The method according to claim 1, wherein the mixture of DNA fragments prepared in step (1) is
- (a) a mixture of DNA fragments in which a modified base or a base is exposed at a cohesive end thereof, obtained by digesting genomic DNA with a restriction enzyme which can digest a DNA regardless of the presence or absence of a modification in a recognition site to generate a cohesive end containing a modified base or a base,
- (b) a mixture of single-stranded DNA fragments or partially single-stranded DNA fragments in which a modified base or a base is exposed in the single-stranded region, obtained by fragmenting genomic DNA and rendering the fragmented genomic DNAs fully or partially single-stranded, or

- (c) a mixture of DNA fragments having a single-stranded region in which a modified base or a base is exposed.
- 3. (Original) The method according to claim 2, wherein the genomic DNA is pretreated with a nuclease capable of digesting a single-stranded DNA, before digesting the genomic DNA with the restriction enzyme to obtain the mixture (a).
- 4. (Original) The method according to claim 2, wherein the genomic DNA or the fragmented genomic DNAs are pretreated with a nuclease capable of digesting a single-stranded DNA, before rendering the fragmented genomic DNAs fully or partially single-stranded to obtain the mixture (b).
- 5. (Currently Amended) The method according to claim 3 [[or 4]], wherein, when the mixture of DNA fragments is brought into contact with the antibody in the step (2), at least one antigen-antibody reaction is performed under conditions in which a monovalent binding is dissociated and a divalent binding is maintained, to separate the mixture into a group consisting of DNA fragments capable of binding to the antibody by a divalent binding, as the group consisting of DNA fragments which form an immunocomplex with the antibody, and another group consisting of DNA fragments capable of binding to the antibody by a monovalent binding, as the group consisting of DNA fragments which do not react with the antibody.
- 6. (Currently Amended) The method according to claim 3 [[or 4]], wherein, when the mixture of DNA fragments is brought into contact with the antibody in the step (2), at least one antigen-antibody reaction is performed under conditions in which a group consisting of DNA fragments showing a high affinity can be separated from another group consisting of DNA fragments showing a low affinity on the basis of the difference between a monovalent binding and a divalent binding, to separate the mixture into a group consisting of DNA fragments capable of binding to the antibody by a divalent binding, as the group consisting of DNA fragments showing a high affinity, and another group consisting of DNA fragments

capable of binding to the antibody by a monovalent binding, as the group consisting of DNA fragments showing a low affinity.

- 7. (Currently Amended) A DNA array obtainable by the method according to any one of claims 1 to 6 and 12 to 13.
- 8. (Original) A group of DNA fragments, characterized by comprising only any one of
- (1) a DNA fragment having cohesive ends containing a modified base or a base at both ends, wherein a modified base is contained in both of the cohesive ends,
- (2) a DNA fragment having cohesive ends containing a modified base or a base at both ends, wherein a modified base is contained in only one of the cohesive ends, or
- (3) a DNA fragment having cohesive ends containing a modified base or a base at both ends, wherein no modified base is contained in both of the cohesive ends.
- 9. (Original) A DNA array characterized in that one or more nucleic acids capable of hybridizing with all or part of DNA fragments contained in the group of DNA fragments of claim 8 are arranged on a substrate.
- 10. (Original) A method of analyzing a modification in a DNA to be assayed, characterized by comprising the steps of:
- (1) preparing a mixture of DNA fragments in which a modified base or a base is exposed, from the DNA to be assayed,
- (2) bringing the mixture of DNA fragments obtained in the step (1) into contact with an antibody specific to the modified base or the base, and separating the mixture into a group consisting of DNA fragments which form an immunocomplex with the antibody and another group consisting of DNA fragments which do not react with the antibody, or a group consisting of DNA fragments showing a high affinity for the antibody and another group consisting of DNA fragments showing a low affinity for the antibody, and
- (3) analyzing all or part of DNA fragments contained in each of the DNA fragment groups with a DNA array.

- 11. (Original) A method of purifying a double-stranded DNA fragment having a cohesive end, characterized by bringing the double-stranded DNA fragment into contact with an antibody specific to a base contained in the cohesive end.
- 12. (New) The method according to claim 4, wherein, when the mixture of DNA fragments is brought into contact with the antibody in the step (2), at least one antigen-antibody reaction is performed under conditions in which a monovalent binding is dissociated and a divalent binding is maintained, to separate the mixture into a group consisting of DNA fragments capable of binding to the antibody by a divalent binding, as the group consisting of DNA fragments which form an immunocomplex with the antibody, and another group consisting of DNA fragments capable of binding to the antibody by a monovalent binding, as the group consisting of DNA fragments which do not react with the antibody.
- 13. (New) The method according to claim 4, wherein, when the mixture of DNA fragments is brought into contact with the antibody in the step (2), at least one antigen-antibody reaction is performed under conditions in which a group consisting of DNA fragments showing a high affinity can be separated from another group consisting of DNA fragments showing a low affinity on the basis of the difference between a monovalent binding and a divalent binding, to separate the mixture into a group consisting of DNA fragments capable of binding to the antibody by a divalent binding, as the group consisting of DNA fragments showing a high affinity, and another group consisting of DNA fragments capable of binding to the antibody by a monovalent binding, as the group consisting of DNA fragments showing a low affinity.